In the Specification

Please amend the specification as shown:

Please delete paragraph [0086] and replace it with the following paragraph:

[0086] To illustrate the method for the identification of motifs of the invention, the example below shows the different matrices constituted in a comparison of motifs performed on a subset of eight sequences based on the reference sequence S V R L G H K D E V **(SEQ ID NO: 1). The peptides that follow are shown in SEQ ID NOS 1-9, respectively, in order of appearance.**

POSITIONS	0 1 2 3 4 5 6 7 8 9
Reference sequence (consensus)	S V R L G H K D E V

Subset of sequences	Alignment
SEQ ID NO. 2 ~~Seq 1~~	S R R L G H K D E V
SEQ ID NO. 3 ~~Seq 2~~	S V R L G H K L E V
SEQ ID NO. 4 ~~Seq 3~~	S R D L G H K D E V
SEQ ID NO. 5 [[Seq4]]	S V R L G H L D V V
SEQ ID NO. 6 ~~Seq 5~~	S V D L G H K T E V
SEQ ID NO. 7 ~~Seq 6~~	S K R L G H K D E V
SEQ ID NO. 8 ~~Seq 7~~	S V R L G H G D G V
SEQ ID NO. 9 ~~Seq 8~~	S V R L G H K S E V

Please delete paragraph [0087] and replace it with the following paragraph:

MUTATION MATRIX A

[0087] Attributed values:

A1 = 0, if motif mutated in relation to the reference sequence

A2 = 1, if another case (motif not mutated in relation to the reference sequence).

POSITION	0 1 2 3 4 5 6 7 8 9
SEQ ID NO. 2 ~~Seq 1~~	1 0 1 1 1 1 1 1 1 1
SEQ ID NO. 3 ~~Seq 2~~	1 1 1 1 1 1 1 0 1 1
SEQ ID NO. 4 ~~Seq 3~~	1 0 0 1 1 1 1 1 1 1
SEQ ID NO. 5 [[Seq 4]]	1 1 1 1 1 1 0 1 0 1
SEQ ID NO. 6 ~~Seq 5~~	1 1 0 1 1 1 1 0 1 1
SEQ ID NO. 7 ~~Seq 6~~	1 0 1 1 1 1 1 1 1 1
SEQ ID NO. 8 ~~Seq 7~~	1 1 1 1 1 1 0 1 0 1
SEQ ID NO. 9 ~~Seq 8~~	1 1 1 1 1 1 1 0 1 1